U.S. Patent Appl. No. 10/009,347 Attorney Docket No. 041322-0284085

### II. REMARKS

By the foregoing amendment, a substitute sequence listing, including a computerreadable form copy and an image form are submitted herewith. The specification is amended to insert the instant sequence listing in place of the previously filed sequence listing. The substitute sequence listing is amended by the addition of feature <223> that indicates that the disclosed sequence is chemically synthesized, as suggested in the sequence listing error report. The applicants also enclose herewith a statement pursuant to 37 C.F.R. § 1.821(f) that the content of the image and computer readable form are the same and do not contain new matter.

Respectfully submitted,
PILLSBURY WINTHROP SHAW PITTMAN LLP

/ thomas a cawley ir /

Thomas A. Cawley, Jr., Ph.D. Registration No. 40944 Telephone: (703) 770.7944

May 10, 2007 CR/LTP P.O. Box 10500 McLean, VA 22102 (703) 905-2000



UNITED STATES DEPARTMENT OF COMMERCE.
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PCL, Sec. 1456
Advantation, Vision. 22313–1480
action. apply.apps

APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO	CONFIRMATION NO
10/009_347	06/26/2002	Leonard C.W. Seymour	P 0284085	3808
	7596 04/19/2007 /INTHROP SHAW PITTI	EXAMINER		
P.O. BOX 10500 MCLEAN, VA 22102			KETTER, JAMES S	
MCLEAN, VA 22102		ART UNIT	PAPER NUMBER	
			(6)th	
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	L	MAIL DATE	DELIVERY MODE	
30 DAYS		04/10/2007	PAPER	

Please find below and/or attached an Office communication concerning this application or proceeding.

If NO period for reply is specified above, the maximum statutory period will apply and will expire 6 MONTHS from the mailing date of this communication.

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FILING DATE

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FIRST NAMED INVENTOR / ATTORNEY DOCKET NO.

EXAMINER

20070329

DATE MARLED:

Please find below and/or attached an Office communication concerning this application or proceeding.

PATENT IN REPYAMMATION

### Commissioner for Patents

The communication filed on 12 January 2007 is non-responsive to the Office communication mailed 12 October 2006 for the reason(s) set forth on the attached Notice to Comply With the Sequence Rules or CRF Diskette Problem Report.

Since the above mentioned reply appears to be bona fide, applicant is given a TIME PERIOD of ONE (I) MONTH or THIRTY DAYS from the mailting date of this notice, which ever is longer, within which to supply the omission or correction in order to avoid abandonment. EXTENSIONS OF THIS TIME PERIOD MAY BE GRANTED UNDER C.F. R. 1.136(a).

The addresses below are effective 5 June 2004. Please direct all replies to the United States Patent and Trademark Office via one (1) of the following:

- Electronically submitted through EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual eFAVE:
- Mailed to
- Mail Stop Sequence Commissioner for Patones

APPLICATION NO.

CONTROL NO

- P.O. Box 22313 1450 Alexandria, VA 22313 1450
- 3. Hand Carry, Federal Express, United Parcel Service or other delivery service to:
- U.S. Patent and Trademark Office
- Mail Stop Sequence Customer Window
- Randolph Building
- 401 Dulancy Street
- Alexandria, VA 22314

Any inquiry concerning this communication should be directed to Examiner Ketter, Art Unit 1636, whose telephone number is (571) 272-4770.

Any industry of a general nature or relating to the status of this application should be directed to the Group recoptionist whose relaptione number is (571) 272-1680.

JAMES KETTER PRIMARY FYATERED

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# NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

18.9	licent must file the items indicated below within the time period set the Office action to which the Notice tlached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the instance of 37 CFR 1 139(a))
The the	nucleotide anti/or amino acid sequence disclosure contained in this application does not comply with requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s).
	<ol> <li>This application clearly fails to comply with the requirements of ST C.F.R. 1.821-1.825. Applicant's attention is directed to the final nulemaking notice published at \$5 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1980). If the effective filling date is on or after July 1, 1998, see the final nulemaking notice published at 63 FR 29820 (June 1, 1998) and 1211 OG 82 (June 23, 1999).</li> </ol>
	<ol> <li>This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).</li> </ol>
	<ol> <li>A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1 821(e).</li> </ol>
	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1 822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
	5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d)
	<ol> <li>The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).</li> </ol>
	7. Other:
	Dicent Must Provide: An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
	An initial or substitute paper copy of the "Sequence Listing", as well as an amendment refrically directing its entry into the application.
appl	A statement that the content of the paper and computer readable copies are the same and, where locable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 5(d).
For	questions regarding compliance to these requirements, please contact:
For	Rules Interpretation, cali (703) 308-4216 or (703) 308-2923 CRF Submission Help, call (703) 308-4212 or 308-2923 entin Software Program Support Technical Assistance
PLE	ASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY

## STIC Biotechnology Systems Branch

# RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/005

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
- APPLICANT, WITH A NOTICE TO COMPLY 61,
  2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A

NOTICE TO COMPLY FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPINCER, TELEPRONE: 571-272-2510, FAX: 571-273-9221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>YERSION 4.4.0 FROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the dishCD-Rom may have been siffered by treatment given to all incoming mail.

Please consider using alternate methods of submission for the dishCD-Rom or replacement dishCD-Rom.

Any teply including a sequence listing in electronic form should NCT be sent to the 2021, zie code soldress for the United States Penert and Trademark Office, and issued should be sent via the following to the indicated soldresses:

- 1. EPS-Bio (<http://www.aspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual - ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Rand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 91/14/95);
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Whodow, Randolph Building, 401 Dalamy Street, Alexandria, VA 27214

Revised 01/19/06

# Baw Sequence Listing Bryor Summary

ERROR DETECTED	SUGGESTED COHRECTION GERIAL NUMBER: 10/009,347
ATTN: NEW BULES CASES	i: Please disregard english "Alpha" headers, which were diserted by PTO Software
IWrapped Nucleics Wrapped Aminos	This number thest at the end of each line "wrapped" down to the next line. This may occur if your file was retired in a word processor after creating it. Please adjust your right inergin to .3; this will prevent "wrapping."
2lovalid Line Length	: The rules require that a line wat exceed 72 characters in length. This includes white spaces.
3Missligned Amino Numbering	The numbering under each $S^A$ amino acid is misaligned. Do not use tab codes between numbers, use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
2Variable Length	Sequenos(s)contain n's or Xua's representing more than one residue. Par Sequence Rielas, each a or Xua cua only represent a stagle residue. Please persent the maximum number of each residue having variable length and indicates in the 2CD>~CZP>~Section that some may be missing.
5Patentin 2.0 "dug"	A "bug" in Patentin version 2.0 has caused the <220><223> section to be missing from amino acid sequences(s)  Mormally, Pinentha would automatically generate this section from the previously coded question acid sequence. Planse removally copy the relevant <220><223> section to the resolution and sequence. This applies to the snandatary <220><223> sections for Artificial or Michaevery aspenses.
Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO.5% (finert SEQ ID NO where "X" is shown) (1) SEQUENCE CHARACTRISTICS; (for no insert any substandings under this heading) (4) SEQUENCE DESCRIPTION:SEQ ID NO.%; (insert SEQ ID NO where "X" is shown) This sequence is intentionally adapted Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
Skipped Sequences (NEW RULES)	Sequence(s)missing. If intentional, please insert the following lines for each skipped sequence. <2100 equence id number <a "chemistrials="" "federal="" "organism"="" "unknown="" (see="" 01="" 06="" 1.523="" 104,="" 1998,="" 29631-322,="" 5"="" 63,="" <2135="" <220-="" <222-="" <2225-="" also="" and="" anterials="" applials="" associated="" by="" explanation.="" feature"="" genetic="" href="https://doi.org/10.000/10.0&lt;/td&gt;&lt;/tr&gt;&lt;tr&gt;&lt;td&gt;Use of n's or Xee's&lt;br&gt;(NEW RULES)&lt;/td&gt;&lt;td&gt;Use of n's and/or Xaa's have been detected in the Sequence Listing.  Per 1 \$25 of Sequence fusiks, use of &lt;220-&lt;225 is MANDATOR') if n's or Xan's are present.  In &lt;200 to &lt;225 section, please explain location of n or Xan, and which residue n or Xan represents.&lt;/td&gt;&lt;/tr&gt;&lt;tr&gt;&lt;td&gt;10lavelid &lt;213&gt;&lt;br&gt;Response&lt;/td&gt;&lt;td&gt;Per 1,823 of Sequence Rules, the only valled &lt;213&gt; responses are: Unissown, Artificial Sequence, or scientific name (Sematispecies). &lt;220&gt;-&lt;223&gt; section is required when &lt;213&gt; response is Unknown or it Artificial Sequence. (see term it before)&lt;/td&gt;&lt;/tr&gt;&lt;tr&gt;&lt;td&gt;11Vie of &lt;2200&lt;/td&gt;&lt;td&gt;Sequence(s) missing the &lt;22/D " identifiers="" if="" is="" jise="" lates"="" mandatory="" no.="" numeric="" of="" off="" or="" please="" pp.="" register,"="" response="" responses.="" sec.="" section="" sequence="" sequence"="" sewie<="" source="" td="" to="" use="" vol.="" white="" wifffeld=""></a>
Patentia 2.0 "hug"	Pleaze do not use "Copy to Drish" function of Pateralin version 2.9. This causes a correspond file, resulting in missing mandatury numeric identifiums and responses (as indicated on now sequence, tilating). Instead, please use "The Manager," or any other manual memors accept file to floggy disk.
U Misse of n/Xaa	"a" can only represent a single <u>mucleotide</u> : "Kan" can only represent a single <u>uning acid</u>
	AMC - STRC Systems Branch - 63/02/06



7 FW 16

RAW SEQUENCE LISTING

PATENT APPLICATION: 05/10/009,347

DATE: 01/17/2007

TIME: 12:24:01

Input Set : %:\efs\01\_17\_07\10009347\_efs\0284085sl.txt Output Ser: N:\CRF4\01172007\3009347.raw

3 <110> AFPLICANT: Seymour, Leonard

4 Fisher, Kerry

5 <120> TITLE OF INVENTION: Modification of Biological Blements

8 <130> FILB REPERENCE: 041322-0284085

10 <140> CUMPERT APPLICATION NUMBER: 10/009.347

11 <141> CURRENT FILING DATE: 2002-06-26

13 <150> PRICE APPLICATION NUMBER: PCT/GB00/02239

14 <151> PRIOR FILING DATE: 2000-06-09

16 <190> PRIOR APPLICATION NUMBER: 99/3359.7 1.7 <151> PRIOR FILING DATE: 1999-06-09

19 <160> NUMBER OF SEQ ID NOS: 1 21 <170> SOFTWARE: Patentin version 3.4

23 <210 > 5EQ ID NO: 1

24 <211> LENGTH: 4

25 <212> TYPE: PRT

26 <213> ORGANISM: Artificial

28 <220> FRATURE:

29 <223 > OTHER INFORMATION: (Artificial sequence 31 <400> SEQUENCE: 1

33 Gly Phe Leu Gly

34 1

LASUFFICIENT OF COMPANY
LASUFFICIENT OF PLEASE GIVE MESORY

SOURCE of genetic material

(See item // on Euro

funcion

fleet)

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: 08/10/009,347 TIME: 12:24:02

DATE: 91/17/2007

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Invalid (211) hesponent | See "<217) Chanism" rasponse is incomplete, per 1.823(h) of New Seebssice Rules. Solid response is Artificial Sequence.

Seq#:1

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/009,347

DATE: 01/17/2007 TIME: 12:34:62

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